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(54) Title: 1 IGAND JAREL CONJUGATES WHICH	H CON	TAIN POLYOXOANIONS OF SULFUR OR PHOSPHORUS		
contain a plurality of chemiluminescent or fluorescent la	abels a	to 100 amino acid residues, bonded to a ligand or receptor, which and a plurality of polyoxoanions of sulfur or phosphorus are useful tvery low nonspecific binding, thereby significantly increasing the		

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Description

Ligand-Lab 1 Conjugat s Which Contain Polyoxoanions of Sulfur or Phosphorus

Technical Field

The present invention relates to novel fluorescentlabeled or chemiluminescent-labeled conjugates for use in specific binding assays for ligands (e.g., antigens and haptens) or receptors (e.g., antibodies, specific binding proteins, and cell surface receptors). The invention further relates to intermediate compounds produced in the synthesis of the novel labeled conjugates, and kits containing such labeled conjugates and/or intermediates.

Background Art

presently, several immunochemical methods exist for the detection of ligands such as haptens, antigens or antibodies. Radioimmunoassay is a widely used method. This method employs a radioisotope-labeled antigen (ligand) to compete with the antigen from a test sample for binding to a specific antibody. U.S. Patent No. 3,709,868 describes such a radioimmunoassay. While by definition, radioimmunoassay is based on the binding of a specific antibody with an antigen or hapten, radioactive binding assays which are based on other specific binding interactions, such as between hormones and their binding proteins, have also been developed.

25 Several non-isotopic immunoassays have been proposed to eliminate the disadvantages associated with radioactive materials. Ligands such as an antigen or hapten and receptors such as an antibody have been labeled with a variety of non-radioactive compounds, including chemiluminescent and fluorescent molecules.

Specific examples of useful chemiluminescent labels are disclosed in G rman OLS No. 2,618,511 and include luminol (3-aminophthalhydrazide or 5-amino-2,3-dihydro-1,4phthalazinedione) and isoluminol (4-

aminophthalhydrazide or 6-amino-2,3-dihydro-1,4-phthalazinedione). The use of N-(4aminobuty1)-N-ethylisoluminol (6-[N-(4-aminobuty1)-Nethylamino]-2,3-dihydrophthalazine-1,4-dione) as a chemiluminescent label is reported in <u>Simpson et al.</u>, <u>Nature</u>, Vol. 279, p. 646

10 (1979). The preparation of chemiluminescent phthalhydrazide labeled ligands is described in U.S. Patent Application Serial No. 927,621, filed July 24, 1978, entitled "Chemiluminescent Phthalhydrazide Labelled Conjugates." U.S. Patent 4,331,808 discloses labeling

ligands with chemiluminescent naphthalene-1,2-dicarboxylic acid hydrazide using straight chain alkyl groups as spacers between the label and the ligand.

While these chemiluminescent compounds have been shown to be suitable labels, they are hydrophobic and,

thereby, increase the hydrophobicity of the ligand complex. For example, column chromatographic purification of synthesized thyroxin-naphthalene-1,2-dicarboxylic acid hydrazide conjugates requires the use of organic solvents (U.S. Patent No. 4,331,808). A documented disadvantage of an increased hydrophobicity is an increase in nonspecific binding effects. This results in a decreased signal to background ratio of immunoassays.

U.S. Patent No. 4,645,646 describes the use of hydrophilic chain-like polymers with recurring functional groups (e.g., proteins) as carriers of multiple chemiluminescent luminol molecules to improve the sensitivity of luminescence immunoassays. However, these

conjugates exhibit non-specific binding properties. In fact, some of the disclosed conjugates xhibit such marked non-specific binding properties to preclude their use in immunoassays (U.S. Patent No. 4,645,646). These examples demonstrate that currently available techniques for coupling chemiluminescent labels to ligands pose an inherent problem. The hydrophobic properties of the chemiluminescent labels decrease the water solubility of the ligands to an extent that the sensitivity of the immunoassay is significantly reduced due to high non-specific background binding.

Labeling of ligands with fluorescent molecules poses very similar problems. Evrain et al., Steroids, vol. 35, 611-619 (1980) describe the synthesis of three

15 fluorescein-labeled derivatives of testosterone using either cysteamine, or 1,3-diaminopropane, or 1,7-diaminoheptane as spacer between the fluorophore and the ligand. All derivatives proved to be highly hydrophobic. For example, analysis of the testosterone

20 fluorescein conjugates by thin-layer chromatography on silica gel required the use of a combination of benzene:ethyl acetate:acetone (1:8:1) or chloroforin:ethanol (7:3) as the solvent system. The hydrophobic nature of this compound is typical of

25 fluorescein-labeled molecules and generally leads to high background readings in immunoassays.

U.S. Patent 4,670,406 describes the use of bifunctional aromatic compounds (e.g., paranitrophenylisocyanate) as rigid coupling compounds for the synthesis of labeled ligands such as fluorescein-labeled digoxin. These rigid coupling reagents are advantageous in that they do not permit the fluorescent

marker to "fold back" onto the ligand, thereby minimizing the possibility of quenching of the fluorescent compound by the ligand. However, as a result of their hydrophobic properties, the rigid coupling reagents further contribute to the loss of the water-solubility of the ligands upon labeling with fluorophores.

U.S. Patent 4,452,886 describes the synthesis of ligand-containing polymers as carriers of multiple photon emitting (fluorescent) compounds. The disclosed polymers 10 include proteins and synthetic or natural polypeptides having a large number of diamino acids for covalent attachment of photon emitting compounds or polymers of such compounds. Similar approaches have also been described in other publications. U.S. Patent 4,604,364 15 discloses tracer compositions for immunoassays which contain photon emitting compounds coupled to ligands via an intermediate support material such as a protein or polypeptide. U.S. Patent Nos. 4,166,105 and 4,169,137 describe antigen detecting reagents which are prepared by 20 covalently linking fluorescent dye molecules to an appropriate antibody through a polymeric backbone having reactive functional groups along the length of its chain. Polymer backbone molecules reported to be suitable are polyethyleneimines (molecular weight range 1200 to 60,000 25 daltons), polypeptides such as polylysines, polyamides such as nylon-6, and low molecular weight (100 to 10,000 daltons) polymeric carboxylic acids. While such polymers help to reduce the loss of antibody binding activity upon fluorescent labeling, they are not suitable to compensate 30 sufficiently for the hydrophobic properties of fluorescent labels such as fluorescein. As a result, such antigen detecting reagents exhibit significant non-specific background binding, thereby limiting the signal to

background ratio of immunoassays. This is clearly demonstrated by a recent observation, in which antibody, molecules labeled with only three fluorescein residues per antibody exhibited an approximately ten-fold higher non-specific binding than the same antibodies labeled with iodine-125 (personal communication, Lisa Shriver-Lake (1990)).

Specific and Non-specific Binding for DTAFand ¹²⁵I-Labeled Goat IgG (Shriver-Lake (1990)).

10

15	Labeled Antigen	Amount of Labeled Antigen Bound to Immobilized anti-Goat IgG	Amount of Labeled Antigen Bound to Immobilized Non-immune IgG	Ratio of Specific to Nonspecific Binding
		(Specific Binding)	(Nonspecific Binding)	
20	125 _{I-labeled} Goat IgG	447.9 ng	30.4 ng	14.73
	DTAF-labele Goat IgG (3.1 DTAF/I		9.52 FU	1.36
25				

FU: arbitrary fluorescence units; DTAF: 5-(4,6-dichlorotriazinyl)aminofluorescein. The specific and nonspecific binding ratios for the DTAF-labeled goat IgG and the 125I-labeled goat IgG were measured using affinity purified anti-goat IgG (Jackson ImmunoResearch, West Grove, PA) covalently attached to cover slips by N-y-maleimidobutyryloxy succinimide ester (GMBS) as described in Bhatia et al, Anal. Biochem., vol. 178, pp. 408-413 (1989).

Thus, even a highly hydrophilic polypeptide chain such as an antibody with a molecular weight of approximately 150,000 apparently cannot compensate for the hydrophobic properties of as few as three fluorophore molecules.

Accordingly, there remains a need for molecular carriers which can link fluorescent or chemiluminescent labels to ligands, such as antigens and haptens, or receptors, such as antibodies, specific binding proteins and cell surface receptors, which do not suffer from the above-mentioned drawbacks. In particular, there remains a need for fluorescent-labeled and chemiluminescent-labeled conjugates which possess good water solubility, and for compounds which contain a plurality of fluorescent or chemiluminescent labels or sites for attaching such labels and which may be conveniently covalently bonded to a ligand, such as an antigen, hapten, or a receptor, such as an antibody, etc., to provide a ligand-label conjugate with good water solubility.

20 Disclosure of the Invention

Accordingly, it is an object of the present invention to provide novel conjugates in which

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chemiluminescent or fluorescent labels are linked to ligands or receptors.

It is another object of the present invention to provide novel conjugates, in which chemiluminescent or fluorescent labels are linked to ligands and receptors, which are hydrophilic.

It is another object of the present invention to provide novel conjugates, in which chemiluminescent or fluorescent labels are linked to ligands and receptors, in which the quenching of the label by the ligand is minimized.

It is another object of the present invention to provide diagnostic kits which contain such conjugates of ligands or receptors and fluorescent or chemiluminescent labels.

It is another object of the present invention to provide novel labeled compounds in which a plurality of fluorescent or chemiluminescent labels are bonded to a linker which possesses a unique functional group for bonding a ligand or a receptor.

It is another object of the present invention to provide novel ligand compounds in which a ligand or a receptor is bonded to a linker molecule which possesses a plurality of functional groups for bonding a plurality of fluorescent or chemiluminescent labels.

These and other objects, which will become apparent during the course of the following detailed description have been achieved by ligand-label conjugates in which an oligopeptide is bonded to a ligand or receptor, and in which at least one of the amino acid residues contains a polyoxoanion group of sulfur or phosphorus and a plurality of the amino acid residues are linked to a chemiluminescent or fluorescent label; the intermediate compounds for coupling the ligand and/or labels to such conjugates; and kits containing such conjugates.

Brief Description of the Drawings

A more complete appreciation of the invention and many of the attendant advantages thereof will be readily obtained as the same become better understood by reference to the following detailed description when considered in connection with the accompanying drawings, wherein:

Figure 1 illustrates the fluorescence lifetime of
the DNP-insulin A-chain-fluorescein conjugate. Shown are
the frequency-dependent phase (D) and modulation (B) data,
together with the best one (- - -) and two (---) component
fits. The precision of the data is approximately ten-fold
better (phase: 0.2 degrees; modulation: 0.002) than
suggested by the size of the data points;

Figure 2 illustrates the relationship between the fluorescence intensity and concentration of the DNP
insulin A-chain-fluorescein conjugate and the DNP-lysinefluorescein conjugate; and

Figure 3 illustrates the binding of different DNP-conjugates to immobilized monoclonal anti-DNP antibody 51.

Best Mode for Carrying Out the Invention

In one aspect, the present invention relates to conjugates in which a natural or synthetic polypeptide oligomer is bonded to a ligand or receptor, and in which at least one of the amino acid residues contains a polyoxoanion of sulfur or phosphorus and a plurality of the amino acid residues are linked to a chemiluminescent or fluorescent label. The polypeptide backbone of the molecule may suitably contain 5 to 100 amino acid

residues, preferably 10 to 50 amino acid residues, most preferably 15 to 25 amino acid residues.

The present conjugates may thus be represented by the general formula (I):

5
$$(aa_1)(aa_2)(aa_3) \dots (aa_n)$$
 (I)

wherein n is an integer of from 5 to 100, preferably 10 to 50, most preferably 15 to 30, and (aa₁) . . . (aa_n) represent amino acid residues, and wherein at least one of the amino acid residues contains a polyoxoanion of sulfur or phosphorus, a plurality of the amino acid residues are linked to a fluorescent or chemiluninescent label, and one of the amino acid residues is bonded to a receptor or ligand.

The present conjugates are easily prepared from

synthetic or natural polypeptide oligomers which contain

three types of functional groups: (i) a unique first

functional group for bonding a ligand or receptor; (ii) at

least one of a second functional group which may be

derivatized to contain a polyoxoanion of phosphorus or

sulfur; and (iii) a plurality of a third functional group

which can be linked to a chemiluminescent or fluorescent

label. Alternatively, the starting oligopeptide may be

one which already contains at least one polyoxoanion, such as insulin A-chain in the tetra-S-sulfonate form and thus, is not required to possess the second type of functional group described above.

is preferably located at one end of the polypeptide oligomer chain. Accordingly, it is preferred that this functional group be either a NH₂- group or a -CO₂H group, since these reactive groups are the natural terminators of polypeptide chains.

In this case, the conjugate may have either formula (II) or (III):

$$H_2N-(aa_1)(aa_2)(aa_3)$$
 $(aa_n)-L$ (II)

$$L-(aa_1)(aa_2)(aa_3)$$
 . . . $(aa_n)-CO_2H$ (III)

15 in which the definitions of formula (I) apply and L is a ligand or receptor.

It is particularly important that the functional group for bonding the ligand be a unique group and that only one ligand is attached to the polypeptide backbone.

20 This is Of particular importance for the detection of

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small molecular weight ligands in competitive
immunoassays. Since the signal in such assays is usually
generated by the release of labeled ligand from the
antibody, increased sensitivity is achieved if the ligand
is bound by only one antibody binding (Fab') site. If two
or more ligands were included in the labeled conjugate,
one ligand could be displaced while another remained bound
to, for example, an antibody-coated solid support.

The term ligand, as used herein, refers to a

molecule such as an antigen or hapten which binds to a
corresponding receptor such as an antibody. Generally,
such ligands include, for example, drugs, hormones,
proteins, vitamins, and infectious agents. The term
receptor, as used herein, includes, for example,
antibodies, specific binding proteins, and cell surface
receptors. As discussed above, the present conjugates
containing ligands and receptors are useful reagents for
quantitative and qualitative measurements of the same
ligand or receptor in homogeneous and heterogeneous
immunoassay systems.

Representative of receptors are antibodies in general, particularly those of the IgG, IgE, IgM and IgA classes, for example hepatitis B antibodies; and representative of ligands are antigenic proteins such as

insulin, chorionic gonadotropin (e.g., HCG),
carcinoembryonic antigen (CEA), myoglobin, hemoglobin,
follicle stimulating hormone, human growth hormone,
thyroid stimulating hormone (TSH), human placental
lactogen, thyroxine binding globulin (TBG), intrinsic
factor, transcobalamin, enzymes such as alkaline
phosphatase and lactic dehydrogenase, hepatitis-associated
antigens such as hepatitis B surface antigen (HB_BAg),
hepatitis r antigen (HB_EAg) and hepatitis core antigen
(HB_CAg) and any antigens derived from HIV such as, e.g.,
gp120. Representative of polypeptide ligands are
angiotensin I and II, C-peptide, oxytocin, vasopressin,
neurophysin, gastrin, secretin, and glucagon.

Since, as peptides, ligands of this general category

comprise numerous available carboxylic acid and amino
groups, coupling to either the terminal -NH₂ or -CO₂H group
of the oligopeptide backbone can proceed according to
conventional peptide condensation reactions such the
carbodiimide reaction, the mixed anhydride reaction, and
so forth, or by the use of conventional bifunctional
reagents capable of coupling carboxylic acid or amino
functions to either the terminal -NH₂ or -CO₂H group of the
oligopeptide backbone. General references concerning the
coupling of proteins to primary amines or carboxylic acids
include <u>Science</u>, vol. 144, p. 1344 (1964); <u>Erlanger et</u>

al., Meth. in Immunology and Immunochemistry, Williams and Chase, eds, Academic, New York, p. 149 (1967); Kopple,

Peptides and Amino Acids, Benjamin, New York (1966); Clin.

Chem., vol. 22, p. 726 (1976); Immunochem., vol. 6, p. 53

(1969) and Lowe et al, Affinity Chromatography, Wiley, New York (1974).

Hapten ligands which themselves contain carboxylic acid functions, and which thereby can be coupled directly to the terminal -NH2 group of the oligopeptide backbone, 10 include the iodothyronine hormones such as thyroxine and liothyronine, as well as other materials such as biotin. valproic acid, folic acid and certain prostaglandins. Representative synthetic routes for preparing carboxylic acid binding analogs of hapten ligands which themselves do 15 not contain an available carboxylic acid function, whereby such analogs can be coupled to the terminal -NH2 group of the oligopeptide backbone by the aforementioned peptide condensation reactions or bifunctional coupling agent reactions, are given in U.S. Patent 4,331,808, which is incorporated herein by reference. Thus, haptens such as carbamazepine, quinidine, digoxin, digitoxin, theophylline, phenobarbital, primidone, diphenylhydantoin, morphine, nicotine, androgens, estrogens, and progesterones may be bonded to the terminal NH2 group of the oligopeptide backbone. Compounds such as cocaine may

also be linked to the terminal -NH₂ group of the oligopeptide backbone. In addition, compounds which possess leaving groups may also be bonded to the terminal NH₂ group. For example, the trinitrophenyl or dinitrophenyl group may be attached by the reaction of fluorotrinitrobenzene or fluorodinitrobenzene with the terminal NH₂ group of the oligopeptide.

As noted above, the ligand or receptor may be bonded to the oligopeptide backbone via a conventional

10 bifunctional spacer. The binding of a low molecular weight hapten, such as, e.g., the trinitrophenyl group, to the oligopeptide via a spacer can be advantageous in that the accessibility of the hapten for binding with its complement (antibody) may be increased. A suitable spacer is, e.g., -(CH₂)_m-S-S- (wherein m is 2 to 10, preferably 4 to 6). For example, the dinitrophenyl group may be bonded to the terminal -NH₂ group of the oligopeptide as shown below.

In the first step S-(2-thiopyridyl)-cysteamine prepared by the reaction of 2,2'-dithiopyridine with cysteamine in methanol (Chong et al, J. Biol. Chem., Vol. 256, pp. 5064-5070 (1981)), is reacted with dinitrofluorobenzene.

5 At the same time, the terminal -NH₂ group of the oligopeptide backbone is derivatized with Traut's reagent (2-iminothiolane) (Jue et al, Biochemistry, Vol. 17, pp. 5399-5406 (1978)). Then, the derivatized oligopeptide is coupled via the free sulfhydryl group to the pyridyldithio group of the derivatized hapten.

Other, bifunctional linkers which may be used to introduce spacers between the ligand or receptor and the oligopeptide backbone include those described in Haugland, Handbook of Fluorescent Probes and Research Chemicals,

Molecular Probes, Inc., Eugene, Oregon (1989) and Pierce Immunotechnology Catalog and Handbook, Pierce, Rockford, Il (1990), e.g., succinimidyl trans-4-(N- maleimidylmethyl)-cyclohexane-1-carboxylate; succinimidyl
4-(p-maleimidylphenyl)butyrate; succinimidyl
6-((iodoacetyl)amino)hexanoate; succinimidyl 6(6(((iodoacetyl)amino)hexanoyl)amino)hexanoate;
5 succinimidyl 3-(2-pyridyldithio)propionate; and
N-succinimidyl S-acetylthioacetate.

Thus, the oligopeptide and the ligand or receptor may be reacted with one of the above-mentioned molecules to introduce one of the following terminal functional

polypeptide-SH
polypeptide-S-S-2-pyridyl

groups:

polypeptide-CCH₂I

polypeptide-maleimidyl

ligand-SH

ligand-S-S-2-pyridyl

O || ligand-CCH₂I ligand-maleimidyl

The derivatized ligand or receptor may then be coupled
with the derivatized oligopeptide as shown in the scheme
below:

Alternatively, either the ligand or oligopeptide can be derivatized with N-succimidyl S-acetylthioacetate resulting in a terminal S-acetyl group which can be deprotected with neutral NH₂OH to yield a free sulfhydryl group which can be coupled as shown above.

Thus, the ligand may be conveniently linked to the oliopeptide via a spacer which contains at least one of the following groups: thioether, dissulfide, and thiosuccinimidyl.

Suitable polyoxoanions have the formula -OMO_n and MO_n, where M may be S or P and n is an integer of from 2 to
3, and include sulfate, sulfonate, sulfinate, phosphate
and phosphonate. These groups are easily introduced into
amino acids and/or the amino acid residues of
polypeptides. For example, sulfate groups may be
introduced into tyrosine by the reaction with sulfuric

acid at low temperatures (Reitz et al, J. Am. Chem. Soc., vol. 68, p. 1024 (1946) and Kohli et al, FEBS Lett., vol. 242, pp. 139-143 (1988)) and may be introduced into tyrosine residues of polypeptides by the use of 5 concentrated sulfuric acid (Ondetti et al, J. Am. Chem. Soc., Vol. 92, pp. 195-199 (1970). Sulfonate groups may be introduced by either the conversion of the free sulfhydryl groups of cysteine residues to S-sulfoderivatives by treatment with sodium tetrathionate, Na2S4O6 10 (Dixon et al, Nature, vol. 188, p. 721 (1960)) or peptides containing disulfide linkages can be reduced with dithiothreitol followed by treatment with sodium tetrathionate (Inglis et al, J. Biol. Chem., vol. 245, p. 112 (1970)). In addition, 3-sulfino-L-alanine is 15 available from Aldrich and may be incorporated in the oligopeptide backbone.

Phosphate groups may be introduced by a variety of methods. Thus, tyrosine may be converted to the O-phosphate derivative by reaction with P₂O₅ in H₃PO₄

20 (Rothberg et al, Proc. Natl. Acad. Sci. USA, vol. 75, pp. 4868-4872 (1978)) or serine may be converted to phosphoserine by treatment with POCl₃ (Neuhaus et al, Biochem. Prepn., vol. 6, p. 75 (1958)). Alternatively, phosphothreonine is commercially available and may be used as a starting amino acid for the production of the

polypeptide oligomer backbone. In addition, histidine can
be derivatized to 7-phosphohistidine by reaction with
phosphoamidate (Hulquist et al, Biochemistry, vol. 5, p.
22 (1966) and Fujitaki et al, Meth. Enzymol., vol. 107,
5 pp. 23-26 (1984)), and arginine can be derivatized to
(ω-N-phosphoarginine by reaction with phosphorus
oxychloride (Thiem et al, Bull. Soc. Chim. Biol., vol. 5,
p. 322 (1962) and Fujitaki et al, Meth. Enzymol., vol. 107,
pp. 23-26 (1984)).

Phosphonate moieties may be introduced into polypeptide oligomers by the reaction of the free sulfhydryl groups of cysteine residues with a phosphonic acid derivative, generated by the reaction of 3-aminopropylphosphonic acid with iodoacetic anhydride

(Meth. Enzymol., vol. 11, p. 532 (1967)).

In addition to the methods described above,

phosphate groups may be introduced into polypeptides by

the enzymatic phosphorylation of amino acid residues in

polypeptides. For example, casein kinase of type II and

camp-dependent kinases phosphorylate serine, while casein

kinases of type I phosphorylate threonine and, to a lesser

extent, serine. Another group of kinases exhibits strict

specificity for tyrosine (Corbin and Hardman, eds., Meth.

Enzymol., vol. 99, Part F, Protein Kinases (1983)).

Similarly, amino acid residues in polypeptides may be enzymatically sulfonated. Thus, tyrosylprotein sulfotransferase catalyzes the sulfonation of proteins at tyrosine residues (Lee et al, J. Biol. Chem., vol. 258, p. 11326 (1983) and Huttner, Meth. Enzymol., vol. 107, pp. 200-233 (1984).

It is preferred that the polyoxoanion is sulfonate.

The polyoxoanion groups may be present as free acids or may be partially or totally neutralized. In the salts formed by partial or total neutralization of the free acid form of the polyoxoanion, the cation may be any suitable ion such as a metal ion or an ammonium ion. Suitable metal ions are those that do not negate the water solubilizing effect of the polyoxoanion and include alkali metals such as lithium, sodium, potassium; alkaline earth metals such as magnesium, calcium, etc.; and transition metals such as, e.g., iron, copper, zinc, etc. Suitable ammonium ions include NH₄⁺ and those in which one or more of the hydrogens have been substituted by an organic

The functional groups which serve to bond the fluorescent or chemiluminescent labels are suitably any of the reactive functional groups found in R of the naturally

occurring amino acids, e.g., -NH₂ in lysine (Lys);
-NHC(NH₂)=NH in arginine (Arg); -CONH₂ in asparagine (Asn)
and glutamine (Gln); -OH in serine (Ser), threonine (Thr),
and tyrosine (Tyr); and -CO₂H in aspartic acid (Asp) and
5 glutamic acid (Glu). It is preferred that the functional
group be one of either NH₂- and -CO₂H, so that one terminus
of the polypeptide backbone may also be conveniently used
as one of the sites for bonding the fluorescent or
chemiluminescent label.

- It is to be understood that the -NH, groups in R of e.g., Lys and the -NH₂ terminal position; the -CONH₂ groups in R of Asn and Gln; the -NHC(NH₂)=NH group in R of Arg; and the imidazole group in R of His are sufficiently dissimilar in reactivity to be classified as different reactive groups for the purposes of the present invention. In other words, the presence of an Asn or Arg residue in the oligopeptide backbone does not prevent the terminal -NH₂ from serving as the unique functional group to bond the ligand or receptor.
- Specific examples of suitable chemiluminescent labels are disclosed in German OLS No. 2,618,511, U.S. Patent 4,331,808, and <u>Haugland</u>, <u>Handbook of Fluorescent Probes and Research Chemicals</u>, Molecular Probes, Inc., Eugene, Oregon (1989) which are incorporated herein by

r fer nc . Thus, suitable labels includ luminol, isoluminol, pyrogallol, lucif rin, and naphthalene-1,2-dicarboxylic acid hydrazide derivatives. These labels may be attached to the oligopeptide backbone by conventional methods. For example, a polypeptide may be labeled with a 7-aminonaphthalene-1,2-dicarboxylic acid hydrazide derivative by condensation of any -CO2H groups on the oligopeptide backbone with a compound of the formula:

as disclosed in U.S. Patent 4,331,808.

those disclosed in <u>Blecka et al</u>, "Immunoassays in Therapeutic Drug Monitoring," <u>Clinics in Laboratory</u>

Medicine, vol. 7, pp. 357-370 (1987) and <u>Haugland</u>,

Handbook of Fluorescent Probes and Research Chemicals,

Molecular Probes, Inc., Eugene, Oregon (1989), which are incorporated herein by reference, and include, e.g.,

fluorescein, rhodamine, anthracene and fluorescamine.

Again, the fluorescent labels may be attached to the

oligopeptide backbone by conventional methods. For example, fluorescein labels may be bonded to a polypeptide which contains a plurality of -CO2H groups by first treating the polypeptide with carbohydrazide, to obtain a carbohydrazide derivative, followed by treating the carbohydrazide derivative with fluorescein isothiocyanate (FITC). This procedure is shown schematically below:

polypeptide-CO₂H + H₂NNHCNHNH₂

polypeptide-CO₂-NHNHCNHNH,

10 polypeptide-CO2NHNHCNHNH2 + FITC

polypeptide-Co20NHNHCNHNHCNH-fluorescein

Similarly, rhodamine X isothiocyanate and 2-anthraceneisothiocyanate may be coupled with the carbohydrazide derivative.

15 Further examples of labels include ruthenium complexes such as those described in U.S. Patent 4,745,076, incorporated herein by reference. These labels

are particularly suitable for time-resolved measurements (U.S. Patent 4,745,076) and electrogenerated chemiluminescent measurements (Eqe et al, Analytical Chem., vol. 56, 2413 (1984) and Zhang et al, J. Phys. 5 Chem., vol. 92, 5566 (1988)).

In regard to the number and distribution of the polyoxoanion groups and the groups for bonding the labels, it is preferred that there be at least two of each of these types of groups. It is especially preferred that 10 there be at least three of both the polyoxoanions and the groups for bonding the label. It is also preferred that the ratio of the number of polyoxoanions to the number of the functional groups for bonding the labels fall within the range of 1:4 to 4:1, more preferably 1:2 to 2:1. 15 the interest of signal strength, it is preferred to keep the number of labels in the molecule relatively high. However, as the ratio of labels to anions in the molecule increases, the hydrophilicity and solubility decrease and the likelihood of signal quenching also increases. 20 this regard, a suitable degree of labeling in the present molecules, and thus a suitable occurrence of groups for bonding the label, is on average about one label per 3 to 10 amino acid residues. A similar degree of substitution is suitable for the polyoxoanion.

It is to be understood that neither the functional groups for bonding the label nor the polyoxoanions need be spaced with strict regularity along the polypeptide backbone. Rather, it is only required that the labels be spaced sufficiently far apart to avoid quenching and that the polyoxoanions be spaced such as to impart the desired hydrophilicity/solubility.

The amino residues contained in the backbone may be any of the naturally occurring α-amino acids represented

10 by the formula H₂N-CH(R)-CO₂H, such as glycine (Gly), alanine (Ala), valine (Val), leucine (Leu), isoleucine (Ile), methionine (Met), proline (Pro), phenylalanine (Phe), tryptophan (Trp) serine (Ser), theronine (Thr), cysteine (Cys), tyrosine (Tyr) asparagine (Asn), glutamine (Gln), aspartic acid (Asp), glutamic acid (Glu), lysine (Lys), arginine (Arg), and histidine (His). The content and sequence of the amino acids are limited only by the need to satisfy the requirement of providing the three types of functional groups discussed above.

For example, if the terminal -NH₂ group of the polypeptide oligomer is to serve as the unique functional group for bonding to the ligand, then the polypeptide oligomer should not contain any amino acid residues which contain -NH₂ groups in R, such as, for example, lysine.

Similarly, when the terminal -CO₂H group is to serve as the unique functional group, the backbone should contain no residues which contain -CO₂H in R. On the other hand, a polypeptide which contains a plurality of aspartic acid or glutamic acid residues will provide a number of -CO₂H groups which can serve to bond the fluorescent or chemiluminescent label.

Thus, the oligopeptide backbone may be either a naturally occurring or synthesized oligopeptide. The

10 synthetic oligopeptide backbone may be synthesized by any conventional procedure. In particular, solid state synthesis utilizing any of the commercially available instruments, such as that produced by Applied Biosystems of Foster City, CA, is suitable. A discussion of the

15 solid state synthesis of oligopeptides is provided in U.S. Patent 3,531,258, which is incorporated herein by reference.

Alternatively, the oligopeptide may be synthesized by recombinant DNA technology. That is, the gene encoding for the desired oligopeptide may be synthesized or isolated and then inserted into a suitable cloning vector which is used to transform a suitable host. The production of oligopeptides by recombinant DNA technology is discussed in U.S. Patents 4,704,362, 4,652,525

4,431,740, 4,440,859, and 4,342,832, which are incorporated herein by reference.

The present conjugates may be assembled by first introducing the polyoxoanion groups into the oligopeptide 5 backbone, followed by bonding the ligand or receptor to the unique functional group and then attaching the labels to the remaining functional groups. Alternatively, the labels may be attached to the backbone before the ligand or receptor is attached or the polyoxoanions may be introduced after either of the other two steps. Thus, in one embodiment, the present invention relates to an intermediate ligand compound which is an oligopeptide, bonded to a ligand or receptor, in which at least one of the amino acid residues has been converted to a 15 polyoxoanion derivative and a plurality of the amino acid residues contain a functional group for bonding a fluorescent or chemiluminescent label. In another embodiment, the present invention relates to an intermediate labeled compound which is an oligopeptide, in 20 which at least one of the amino acid residues has been converted to a polyoxoanion derivative, a plurality of the amino acid residues are linked to a fluorescent or chemiluminescent label, and having a unique functional group for bonding a ligand or receptor.

These intermediate compounds may be represented by the general formulae

$$L-(aa_1)(aa_2)(aa_3) (aa_n)-CO_2H$$
 (IV)

$$H_2N-(aa_1)(aa_2)(aa_3)$$
 $(aa_n)-L$ (V)

wherein L and n are as defined above and at least one of the amino acid residues contains a polyoxoanion of phosphorus or sulfur and a plurality of the amino acid residues have a functional group for bonding a chemiluminescent or fluorescent label; or

10
$$H_2N-(aa_1)(aa_2)(aa_3) \dots (aa_n)-CO_2H$$
 (VI)

wherein n is as defined above and at least one of the amino acid residues contains a polyoxoanion of phosphorus or sulfur and a plurality of the amino acid residues are bonded to a chemiluminescent or fluorescent label.

the insulin A-chain and fragments of cholecystokininpancreozymin which may be sulfated as disclosed in <u>Ondetti</u>
et al, <u>J. Am. Chem. Soc.</u>, Vol. 92, pp. 195-199 (1970) some
of which are shown below.

SO₃H | H₂N-Phe-Asp-Met-Trp-Gly-Met-Tyr-Asp-CO₂H

SO₃H | H₂N-Phe-Asp-Met-Trp-Gly-Met-Tyr-Asp-Arg-Asp-CO₂H

A preferred embodiment of the oligopeptide backbone of present conjugates is represented by insulin A-chain or 5 a polypeptide having substantially the same number of amino acid residues and molecular weight as insulin A-chain, with one terminal amino or carboxy group for covalent attachment of a ligand or receptor molecule, at least two functional groups for covalent attachment of a label at a distance from the ligand or receptor site 10 sufficient to minimize quenching of the labels by the ligand or receptor, and at least two functional groups which may be converted to a polyoxoanion of phosphorus or sulfur. Such oligopeptides include insulin A-chain; oligopeptides having substantially the same number of amino acid residues as insulin A-chain, at least two cysteine residues and at least one amino acid residue having a -CO₂H group in R, such as Glu or Asp, but only one -NH2 group; and oligopeptides having substantially the 20 same number of amino acid residues as insulin A-chain, at least two cysteine residues, and at least one amino acid

residue having a $-NH_2$ group in R, such as Lys, but only one $-CO_2H$ group. Examples of such peptides may be represented by the formula

$$H_2N-(aa_1)(aa_2)(aa_3)$$
 . . . $(aa_n)-CO_2H$ (VII)

in which n is about 21, at least two of the amino acid residues are cysteine, at least one of the amino acid residues has a -CO₂H group in R, such as Glu or Asp, and none of the remaining amino acid residues have -NH₂ groups in R, such as Lys; or n is defined as above, at least two of the amino acid residues are cysteine, at least one of the amino acid residues has a -NH₂ group in R, such as Lys, and none of the remaining amino acid residues contain a -CO₂H group in R, such as Glu or Asp.

The A-chain of insulin represents an ideal backbone

15 for the synthesis of a sulfonated oligomeric carrier

molecule. The A-chain of insulin is a hydrophilic peptide

containing 21 amino acid residues and has a molecular

weight of approximately 2500 daltons.

Four of the amino acids are cysteine residues with a free sulfhydryl group which can easily be derivatized to S-sulfonates, <u>J. Am. Chem. Soc.</u>, vol. 88, 5625-5635 (1966). Two carboxyl groups from glutamate residues 4 and

17 and one terminal carboxyl group from an asparagine residue provide three sites for attachment of fluorescent or chemiluminescent molecules. The insulin A-chain contains one terminal amino group for covalent attachment of a single ligand or receptor molecule. Since the sites for attachment of the fluorescent or chemiluminescent labels are 4, 17, and 21 amino acid residues away from the ligand or receptor attachment site, the possibility of quenching of the labels by the ligand or receptor is minimized.

Thus, insulin A-chain in the tetra-S-sulfonate form may be easily coupled to an antigen, such as the dinitrophenyl group, via the terminal -NH2 group by reaction with 1-fluoro2,4-dinitrobenzene, to obtain dinitrophenyl-insulin A-chain (DNP-insulin A-chain).

Derivatization of the carboxyl groups of DNP-insulin A-chain with carbohydrazide, followed by reaction with fluorescein isothiocyanate (FITC) gives DNPinsulin A-chain-fluorescein (DNP-Ins-Fl), the structure of which is shown below:

The ligand-insulin A-chain-fluorescein conjugates
(Ligand-Ins-Fl) are ideally suited for use in assays for
the free ligand. For example, the amount of non-specific
binding of DNP-Ins-Fl to immobilized anti-DNP IgG

(antibody #51) is only about one-third that of a conjugate
in which a dinitrophenyl group is linked to fluorescein
via a lysine residue, DNP-Lys-fluorescein (DNP-Lys-Fl)
(see Table II). Further, the amount of DNP-Ins-Fl
specifically bound to antibody #51 is about 1.7-fold
higher than that of DNP-Lys-Fl.

In Figure 3 the binding of the different DNP conjugates to anti-DNP-Sepharose at a 3:1 molar ratio of conjugate to immobilized antibody is shown. After extensive washing of the matrix, a 1.6-fold difference in binding of DNP-Lys-Fl (62.7%) versus DNP-Ins-Fl (38.8%)

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was observed. In a attempt to identify the factors contributing to this significant difference, the binding of DNP-Ins without attached fluorescein molecules was also determined. The data in Table II reveal an almost identical extent of binding of DNP-Ins and DNP-Ins-Fl to anti-DNP-Sepharose, suggesting that the increased number of fluorophores in the DNP-Ins-Fl conjugate compared to the DNP-Lys-Fl conjugate is not a critical factor.

In order to determine whether the differences in the 10 extent of binding shown in Figure 3 are a result of a different affinity of the immobilized anti-DNP antibody for both conjugates, DNP-Ins-Fl and DNP-Lys-Fl, the displacement of the different DNP conjugates from the antigen binding sites of immobilized anti-DNP antibody #51 15 by DNP-lysine was determined. The analyses revealed that under comparable conditions, the displacement of DNP-Ins-Fl is much more efficient than that of DNP-Lys-Fl (see Table III). However, the data in Table III also demonstrate that the displacement of DNP-Ins-Fl is two-20 fold more efficient than that of DNP-Ins, whereas the extent of binding of DNP-Ins-Fl and DNP-Ins was almost identical (Table II). Therefore, a difference in affinity cannot be the only factor responsible for the different binding efficiencies of the DNP conjugates to antiDNP-Sepharose.

Table II compares the extent of binding of the different DNP conjugates to immobilized anti-DNP IgG (antibody #51) with that to immobilized non-immune IgG. The data show that the extent of non-specific binding of 5 DNP-Lys-Fl to immobilized non-immune IgG is three-fold higher than that of DNP-Ins-Fl. After subtraction of nonspecific binding to non-immune IgGSepharose from binding to anti-DNP-Sepharose, a fundamentally different binding pattern becomes evident. The amount of DNPIns-Fl that 10 specifically is bound to the antigen binding sites of immobilized antibody #51 (0.7 mole conjugate/mole antibody) is 1.7-fold higher than that of DNP-Lys-Fl (0.4 mole conjugate/mole antibody). These results show that the different extent of binding of DNP-Lys-Fl and DNP-Ins-15 Fl to anti-DNP-Sepharose given in Figures 3 is a consequence of the high nonspecific binding of DNP-Lys-Fl.

In another aspect, the present invention relates to kits which contain the present conjugates and/or intermediate compounds. Such kits may take any form that is suitable for the employment of the present conjugates and/or intermediate compounds in an assay for a ligand and/or receptor. In particular, the present kits may contain a vial or other container which may contain the present conjugate and/or intermediate compound in solution

or powdered form. If the present conjugate and/or intermediate compound is in solution form, the solution may also contain other materials, such as, e.g., a preservative, buffer, salt, etc. The kit may also contain a standard sample of either the ligand or receptor of the present conjugate or the binding complement of the ligand or receptor. Such standard samples may be in the form of a solution of known concentration and/or activity or a sample of known weight and/or activity. In addition, the present kits may contain samples in which the binding complement of the ligand or receptor of the conjugate is immobilized on a solid support, such as a glass slide, glass beads, gelatin beads, polymer matrix, etc.

The present conjugates and kits are useful for assaying ligands or receptors and/or the binding complements of the ligands or receptors, in specific binding assays in which the displacement or competitive binding of the present conjugates are used to determine the amount or concentration of the free ligand or receptor (or complement thereof) present in a sample. The sample may be of biological or nonbiological origin and may be originally obtained in gaseous, liquid or solid form.

Other features of the invention will become apparent in the course of the following descriptions of exemplary

embodiments which are given for illustration of the invention and are not intended to be limiting thereof.

General Methods

Fluorescence intensity was measured on a SLM 8000 5 fluorimeter at 490 nm excitation and 519 nm emission. Individual fluorescence standard curves with known concentrations of the different DNP-fluorescein conjugates were used to calculate the amounts of DNP-conjugates from fluorescence units. Antibodies were coupled to tresyl 10 chloride-activated Sepharose 4B (Sigma) at a concentration of 200 ug antibody/100 mg gel according to the manufacturer's recommendation. The amount of immobilized antibody was determined using a Coomassie blue assay as described in Ahmad et al, Anal. Biochem., vol. 148, pp. 15 533-541 (1985). Peptide and protein determinations in fluid-phase were performed by UV absorbance at 280 nm, by the bicinchoninic acid method (Smith et al, Anal. Biochem., vol. 150, pp. 76-85 (1985)) using the Pierce BCA Protein Assay Reagent (Pierce), and by the FolinCiocalteau 20 method (Lowry et al, J. Biol. Chem., vol. 193, pp. 265 (1951)). The A-chain of bovine insulin (oxidized to the S-sulfonate form) (Sigma), the DNP-Ins conjugate, and the DNPIns-Flu conjugate were radiolabeled with Na¹²⁵I

(Amersham, Arlington Heights, IL) using immobilized

chloramine-T (IodoBeads; Pierce, Rockford, IL) (<u>Petrella</u> et al, <u>J. Immunol. Methods</u>, vol. 104, pp. 159-172 (1987)).

Synthesis of the DNP-Insulin A-Chain-Fluorescein Conjugate

The synthesis was performed in three steps: 1)

5 coupling of 1-fluoro-2,4-dinitrobenzene (FDNB) to the terminal amino group of the insulin A-chain oxidized to the tetra-S-sulfonate form, 2) derivatization of the carboxyl groups with carbohydrazide, and 3) coupling of fluorescein isothiocyanate (FITC) to the hydrazide

10 derivative of DNP-insulin A-chain.

Step 1. Insulin A-chain in the tetra-S-sulfonate form (Sigma, St. Louis, MO) (25 mg; approximately 8 μmoles), mixed with ¹²⁵I-labeled S-sulfonated insulin A-chain (26 nmole, 5x10⁷ cpm), was incubated with 50 μmol (9.3 mg) of FDNB (Aldrich, Milwaukee, WI) in a total volume of 6.4 mL containing 100 mM NaHCO₃ and 19% (v/v) ethanol. After two hours of incubation at 37°C, the reaction mixture was subjected to Sephadex G-10 size exclusion chromatography to remove noncoupled FDNB molecules. The peptide fractions, eluted with 19% (v/v) ethanol in H₂O, were lyophilized and stored at -20°C. The ratio of coupled FDNB residues per insulin A-chain was calculated from the specific radioactivity of the peptide

and the UV absorbance at 365 nm using an extinction coefficient of 1.6 x 104 M⁻¹ cm⁻¹ (Hirs, Meth. Enzymol., XI, pp. 548-555 (1967)). Preparations containing less than one DNP residue per insulin A-chain were subjected to a second derivatization with freshly added FDNB as described above.

Step 2. DNP-derivatized insulin A-chain (tetra-S-sulfonate form) (DNP-Ins) (3.5 mg; approximately 1.2 μmol) was dissolved in 800 μl of 10 mM sodium phosphate-150 mM

10 NaCl, pH 7.4. After the addition of 165 mg (1.8 mmol) of carbohydrazide (Aldrich) and 20 mg (0.1 mmol) of 1-ethyl-3-(3-dimethylaminopropyl)-carbodiimide (EDC) (Sigma), the pH was adjusted to pH 5.0 with hydrochloric acid. The reaction mixture (total volume 950 μl) was incubated at 15 room temperature for 18 hours (the pH was maintained at 5.0) and then subjected to Sephadex G-10 size exclusion chromatography. The peptide fraction, eluted with 19% (v/v) ethanol in H₂O, was lyophilized and stored at -20°C.

Step 3. Carbohydrazide-derivatized DNP-insulin

20. A-chain (tretra-S-sulfonate form) (0.35 mg; approximately
0.1 nmol) was dissolved in 950 μl of 50 mM NaHCO₃

containing 19% (v/v) ethanol. FITC (Aldrich) (16 mg; 41

μmol) was added, and after 18 hours of incubation at room
temperature in the dark, the reaction mixture was

subjected to Sephadex G-10 size exclusion chromatography. The peptide fraction, eluted with 19% (v/v) ethanol in H₂O, was lyophilized, redissolved in 50 mM NaHCO₃ containing 19% (v/v) ethanol, and rechromatographed on Sephadex G-10 to remove traces of any non-covalently bound fluorophores. The number of covalently attached fluorescein molecules was spectrophotometrically determined at 494 nm using an extinction coefficient of 7.6 x 10⁴ M⁻¹cm⁻¹ (Wilderspin et al, Anal. Biochem., vol. 132, pp. 449-455 (1983)).

Spectrophotometric analyses of the purified DNPinsulinA-chain-fluorescein conjugate (DNP-Ins-Fl)
confirmed the theoretical ratio of one DNP residue and
three fluorescein residues per insulin A-chain molecule as
shown below in Table I.

Table I. Molar ratio of covalently attached FDNB and FITC molecules per insulin A-chain.

20	Derivatization of A-chain insulin A-ch with:		residue/insulin
		Expected	Found

		Expected	Found
	FDNB	1.0	1.1 ± 0.1
25	FITC	3.0	3.2 ± 0.1

synthesis of the DNP-Lysin -Pluorescein Conjugate

ε-DNP-L-lysine hydrochloride (Research Organics Inc., Cleveland, Ohio 44125) (10 mg; 0.029 mmol) was dissolved in 1.0 ml of 0.5 M NaHCO3 and adjusted to pH 9.3 5 with NaOH. Pluorescein isothiocyanate (Aldrich) (10 mg; 0.026 mmol) was added, and the reaction mixture (total volume 1.2 ml) was incubated in the dark for 30 min at room temperature. Purification of the synthesized DNPlysine-fluorescein conjugate (DNP-Lys-Pl) was performed by . 10 reverse phase HPLC on an Ultrasphere ODS C-18 column (Beckman, CA) at a flow rate of 0.75 ml/min using a linear gradient from 5% (v/v) methanol (in H2O) to 100% methanol (within 15 min). The DNP-Lys-Pl conjugate fractions from several chromatographic runs were combined, lyophilized, 15 and stored at -20°C. Rechromatography of an aliquot of the combined fractions under identical conditions demonstrated 95% purity of the DNP-Lys-Fl conjugate. structure of DNP-Lys-Fl is shown below.

Determination of Pluoresc nce Lif time

The fluorescence lifetime of the DNP-fluorescein conjugates was measured on an ISS Greg-200 variable frequency phase fluorimeter as described previously (Lakowicz et al, Biophys. J., vol. 46, pp. 463-477 (1984) and Gratton et al, Biophys. J., vol. 46, pp. 479-486 (1984)) using 1-chloro-bis-(phenylethynyl)anthracene (Aldrich) in ethanol (3.791 ns) as a reference (Thompson et al, Anal. Chem., vol. 60, pp. 670-674 (1988)). The 10 resulting phase and modulation data for DNP-Ins-FL were fit to one, two, and three emissive components using a Simplex algorithm supplied by ISS. The results are shown in Figure 1. The data fitted best $(X^2=1.26)$ to two components: the first comprises about two thirds of the 15 mixture and exhibited a lifetime corresponding closely to that of fluorescein $(4.091 \pm 0.177 \text{ ns}; 64 \pm 7\%)$, and a shorter lifetime component comprising about one third of the mixture (2.068 ± 0.150 ns; 36%). Fitting these data to one component gave an unacceptable X^2 ($X^2 = 33$) and non-20 random differences between the data and calculated values. When the data were fit to a three component model, the third component was weak (<5%), poorly defined (1.1 ± 3.2 ns), and gave no improvement in X^2 . Therefore, a third component appears to be unnecessary to fit the data. 25 Since the DNP moiety has the ability to quench the

fluorescein emission, the data are best explained by two fluorescein residues being remote from and unquenched by the DNP group with the third fluorescein residue being closer and subject to quenching. In accordance with this explanation is the observation that the fluorescence of the DNP-lysine-fluorescein derivative was substantially quenched as shown by its reduced quantum efficiency and subanosecond fluorescence lifetime.

In addition, the relationship between the

10 fluorescence intensity and concentration for DNP-Ins-Fl

and DNP-Lys-Fl is given in Figure 2.

Specific and Nonspecific Binding of DNP-Conjugates

Specific binding of the DNP-conjugates was
determined with the murine monoclonal anti-DNP antibody

15 #51 (Stanley et al, J. Immunol. Methods, vol. 64, pp.
119-132 (1983) and Steward et al, J. Immunol. Methods,
vol. 78, pp. 173-190 (1985)) coupled to tresyl chlorideactivated Sepharose 4B (anti-DNP-Sepharose). The
monoclonal IgGl antibody #51 (purified from ascites) was

20 kindly provided by Dr. M.W. Steward (Dept. of Medical
Microbiology, London School of Hygiene and Tropical
Medicine, London, U.K.). Nonspecific binding was measured
with Sepharose 4B-immobilized non-immune mouse IgG

purified from mouse serum (Jackson ImmunoResearch Laboratories, West Grove, PA) (IgG-Sepharose). After removal of buffer by vacuum filtration on a scintered glass funnel, 50 mg aliquots of Sepharose 4B-antibody 5 matrices were transferred to plastic conical vials and incubated with 125I-labeled DNPinsulin A-chain-fluorescein conjugate (1.6 x 106 cpm/nmole), or 125I-labeled DNPinsulin A-chain conjugate without attached fluorescein residues (3.3 x 106 cpm/nmole), or non-radiolabeled DNP-10 Lys-Fl at a 3 to 1 molar ratio of DNP-derivative to immobilized IgG in 10 mM phosphate/150 mM NaCl, pH 7.6, overnight at 4°C on a rocking platform. Unbound conjugate was recovered by poking a hole in the bottom of the plastic vial with a 30 gauge needle, loading the vial on 15 the top of a glass test tube, and centrifuging. Subsequent washing steps were performed with 150 μ l of 10

Subsequent washing steps were performed with 150 μ l of 10 mM phosphate/150 mM NaCl/0.1% (v/v) Triton X-100, pH 7.6. Aliquots of the solutions collected in the bottom of the test tubes after centrifugation were assayed for

20 radioactivity or fluorescence. The results are shown in Table II and Figure 3.

Table II. Specific and nonspecific binding of DNP-conjugates.

5	DNP- Conjugate	DNP- Conjugate Applied	Ratio of Applied DNP- Conjugate to Immobilized IgG	DNP- Conjugate Bound	Ratio of Bound DNP- Conjugate to Immobilized IgG
		(nmoles)	(moles/ moles)	(nmoles)	(moles/ moles)
15	I. Binding to anti-DN IgGl			·	
	DNP-Lys-Fi	0.59	3.0	0.37±0.02	1.9
	DNP-Ins-F1	L 0.29	3.0	0.11±0.03	1.2
	DNP-Ins	0.29	3.0	0.11±0.01	1.2
20	II. Bindi to non- immune Igo	_			
	DNP-Lys-F	2.60	3.0	1.27±0.07	1.5
2.5	DNP-Ins-F	2.53	3.0	0.44±0.16	0.5

Specificity and Effectivity of Displacement of DNP-Conjugates

Aliquots of anti-DNP-Sepharose (50 mg) were incubated with ¹²⁵I-labeled DNP-insulin A-chain, or ¹²⁵I
30 labeled DNPinsulin A-chain-fluorescein, or DNP-lysine-fluorescein and washed with 10 mM phosphate/150 mM

15

NaCl/0.1% (v/v) Triton X100, pH 7.6, as described above in Specific and Nonspecific Binding of DNP-Conjugates. When the wash solutions showed background levels of radioactivity or fluorescence, 150 µl of 10 mM

5 phosphate/150 mM NaCl/0.1% (v/v) Triton X-100, pH 7.6, containing the compound to be tested for displacement of antibody-bound DNP-conjugate was added to the anti-DNP-Sepharose aliquots, and centrifuged as described.

Displacement was determined by assaying aliquots of the solutions collected in the bottom of the test tubes after centrifugation for radioactivity or fluorescence. The results are shown in Table III.

Table III. Displacement of DNP-conjugates from immobilized monoclonal anti-DNP antibody by DNP-lysine.

20		Ratio of Free DNP- Lysine to Antibody- Bound DNP-Conjugate	Displacement of Antibody-Bound DNP- Conjugate by DNP- Lysine	
		[mole/mole]	[pmole ± S.D.]	
	DNP-Lys-Fl	2.0	0.45 ± 0.14	
	DNP-Ins-Fl	3.4	3.20 ± 0.70	
25	DNP-Ins	3.4	1.56 ± 0.01	

Obviously, numerous modifications and variations of the present invention are possible in light of the above teachings. It is therefore to be understood that within the scope of the appended claims, the invention may be practiced otherwise than as specifically described herein.

CLAIMS

- A ligand-label conjugate, which is an oligopeptide of 5 to 100 amino acid residues, which is bonded to a ligand or receptor, wherein at least one of said amino acid residues contains a polyoxoanion of sulfur or phosphorus and a plurality of said amino acid residues are linked to a chemiluminescent or fluorescent label.
 - 2. The conjugate of Claim 1, wherein the number of amino acid residues in said oligopeptide is from 10 to 50.
- 3. The conjugate of Claim 1, wherein the number of 10 amino acid residues in said oligopeptide is from 15 to 25.
- 4. The conjugate of Claim 1, wherein said label is selected from the group consisting of luminol, isoluminol, pyrogallol, luciferin, naphthalene-1,2-dicarboxylic acid hydrazide derivatives, fluorescein, rhodamine, anthracene, fluorescamine, and ruthenium complexes.
 - 5. The conjugate of Claim 1, wherein said polyoxoanion of sulfur or phosphorus is selected from the group consisting of sulfate, sulfonate, sulfinate, phosphate and phosphonate.

- 6. The conjugate of Claim 1, wherein said polyoxoanion of sulfur or phosphorus is sulfonate.
- 7. The conjugate of Claim 1, wherein said oligopeptide has substantially the same amino acid residues and molecular weight as insulin A-chain, said amino acid residues containing a polyoxoanion of sulfur or phosphorus are S-sulfonate-Cys, said amino acid residues linked to said labels are Glu or Asp and the CO₂H-terminal amino acid residue, and said ligand is linked to the H₂N-terminal amino acid residue.
 - 8. The conjugate of Claim 1, wherein said oligopeptide has the same amino acid sequence as insulin A-chain.
 - 9. The conjugate of Claim 1, which has the formula:

- 10. The conjugate of Claim 1, wherein said ligand or receptor is bonded to said oligopeptide via a spacer group.
- 11. The conjugate of Claim 10, wherein said spacer 5 group contains a thioether group, a disulfide group, or a thiosuccinimidyl group.
 - 12. A ligand compound which is an oligopeptide of 5 to 100 amino acid residues which is bonded to a ligand or receptor, wherein at least one of said amino acid residues contains a polyoxoanion of phosphorus or sulfur and a plurality of said amino acid residues contain a functional group for bonding a chemiluminescent or fluorescent label.
- 13. The ligand compound of Claim 12, wherein the number of amino acid residues in said oligopeptide is from15 10 to 50.
 - 14. The ligand compound of Claim 12, wherein the number of amino acid residues in said oligopeptide is from 15 to 25.
- 15. The ligand compound of Claim 12, wherein said20 polyoxoanion of sulfur or phosphorus is selected from the

group c nsisting of sulfat , sulfonate, sulfinate, phosphat and phosphonat .

- 16. The ligand compound of Claim 12, wherein said polyoxoanion of sulfur or phosphorus is sulfonate.
- oligopeptide has substantially the same amino acid residues and molecular weight as insulin A-chain, said amino acid residues containing a polyoxoanion of sulfur or phosphorus are S-sulfonate-Cys, said amino acid residues

 linked to said labels are Glu or Asp and the CO2H-terminal residue, and said ligand is linked to said H2N-terminal amino acid residue.
- 18. The ligand compound of Claim 12, wherein said oligopeptide has the same amino acid sequence as insulin 15 Achain.
 - 19. The ligand compound of Claim 12, which has the formula:

- 20. The ligand compound of Claim 12, wherein said functional group for bonding a chemiluminescent or fluorescent label is -NH, or -CO,H.
- 21. The ligand compound of Claim 12, wherein said
 5 ligand or receptor is bonded to said oligopeptide via a spacer group.
 - 22. The ligand compound of Claim 21, wherein said spacer group contains a thioether group, a disulfide group, or a thiosuccinimidyl group.
- 23. A labeled compound which is an oligopeptide of 5 to 100 amino acid residues in which at least one of said amino acid residues contains a polyoxoanion and a plurality of said amino acid residues are bonded to a chemiluminescent or fluorescent label and one of said 15 amino acid residues contains a unique functional group for bonding a ligand or receptor.
 - 24. The labeled compound of Claim 23, wherein the number of amino acid residues in said oligopeptide is from 10 to 50.

- 25. The label d compound of Claim 23, wherein the number of amino acid residues in said oligopeptide is from 15 to 25.
- 26. The labeled compound of Claim 23, wherein said chemiluminescent label is selected from the group consisting of luminol, isoluminol, pyrogallol, luciferin, naphthalene-1,2-dicarboxylic acid hydrazide derivatives, fluorescein, rhodamine, anthracene, fluorescamine, and ruthenium complexes.
- 27. The labeled compound of Claim 23, wherein said polyoxoanion of sulfur or phosphorus is selected from the group consisting of sulfate, sulfonate, sulfinate, phosphate and phosphonate.
- 28. The labeled compound of Claim 23, wherein said polyoxoanion of sulfur or phosphorus is sulfonate.
 - 29. The labeled compound of Claim 23, wherein said unique functional group for bonding a ligand or receptor is -NH₂ or -CO₂H.
- 30. The labeled compound of Claim 23, wherein said oligopeptide has substantially the same amino acid residues and molecular weight as insulin A-chain, said

amino acid residu s containing a polyoxoanion of sulfur or phosph rus are S-sulfonat -Cys, said amino acid residues linked to said labels are Glu or Asp and the CO₂H-terminal residue, and said ligand is linked to the H₂N-terminal amino acid residue.

- 31. The labeled compound of Claim 23, wherein said oligopeptide has the same amino acid sequence as insulin Achain.
- 32. The labeled compound of Claim 23, which has the

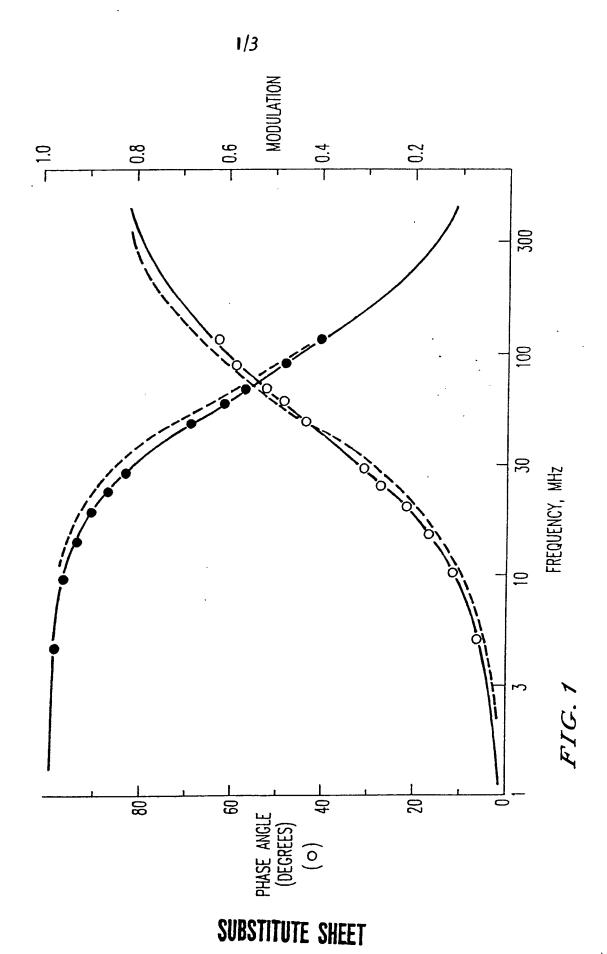
33. A kit, comprising (i) a ligand-label conjugate, which is an oligopeptide of 5 to 100 amino acid residues, which is bonded to a ligand or a receptor, wherein at

least on of said amino acid residues contains a polyoxoanion of sulfur or phosphorus and a plurality of said amino acid residues are linked to a chemiluminescent or fluorescent label, and (ii) a binding complement of said ligand or receptor.

- 34. The kit of Claim 33, wherein the number of amino acid residues in said oligopeptide is from 10 to 50.
- 35. The kit of Claim 33, wherein the number of 10 amino acid residues in said oligopeptide is from 15 to 25.
- 36. The kit of Claim 33, wherein said chemiluminescent label is selected from the group consisting of luminol, isoluminol, pyrogallol, luciferin, naphthalene-1,2-dicarboxylic acid hydrazide derivatives, fluorescein, rhodamine, anthracene, fluorescamine, and ruthenium complexes.
- 37. The kit of Claim 33, wherein said polyoxoanion of sulfur or phosphorus is selected from the group20 consisting of sulfate, sulfonate, sulfinate, phosphorate and phosphonate.
 - 38. The kit of Claim 33, wherein said polyoxoanion of sulfur or phosphorus is sulfonate.

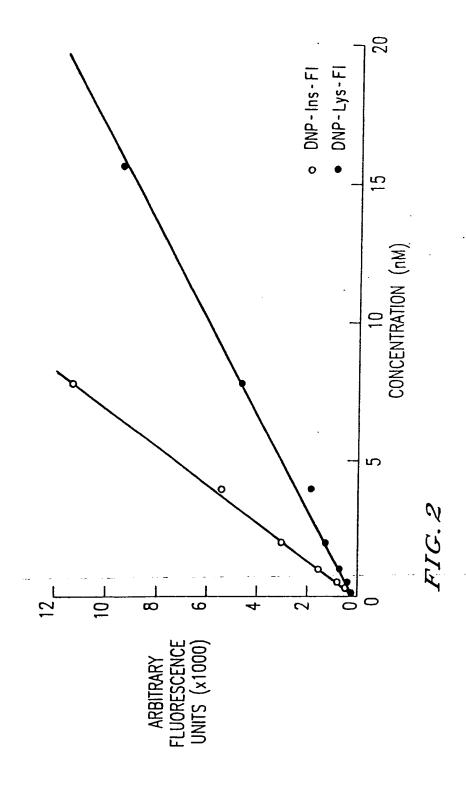
- 39. The kit of Claim 33, wh rein said oligopeptide has substantially the sam amino acid residues and molecular weight as insulin A-chain, said amino acid residues containing a polyoxoanion of sulfur or phosphorus are S-sulfonate-Cys, said amino acid residues linked to the labels are Glu or Asp and the CO2H-terminal residue, and said ligand is linked to the H2N-terminal amino acid residue.
- 40. The kit of Claim 33, wherein said oligopeptide 10 has the same amino acid sequence as insulin A-chain.
 - 41. The kit of Claim 33, wherein said ligand-label conjugate has the formula:

- 42. The kit of Claim 33, further comprising a sample of said ligand or receptor.
- 43. The kit of Claim 42, wherein said sample of said ligand or receptor contains a known amount of said ligand or receptor.
 - 44. The kit of Claim 33, wherein said binding complement of said ligand or receptor is present in the form of a sample of a known amount.
- 45. The kit of Claim 33, wherein said ligand or 10 receptor is bonded to said oligopeptide via a spacer group.
 - 46. The kit of Claim 45, wherein said spacer group contains a thioether group, a disulfide group, or a thiosuccinimidyl group.

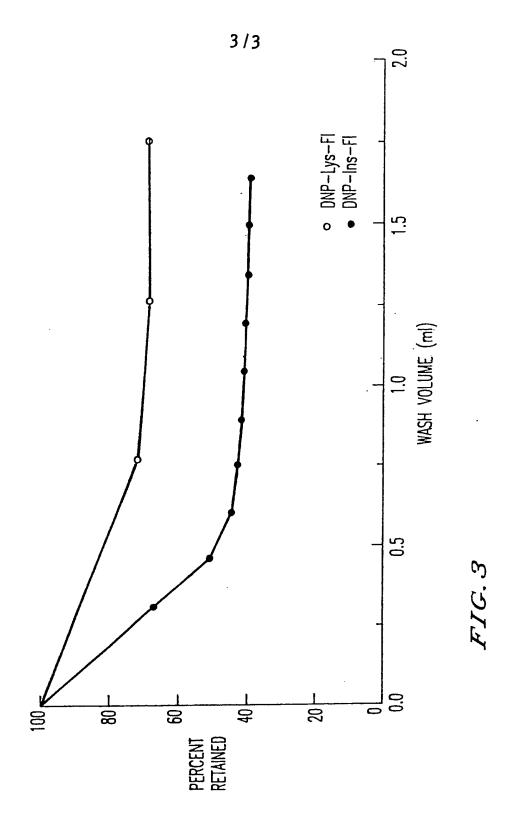


WO 91/16344 PCT/US91/02212





SUBSTITUTE SHEET



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INTERNATIONAL SEARCH REPORT

International Antibication to PCT/US91/02212

1. CLASSIFICATION OF SUBJECT MATTER of several classification symbols apply, imprate all 5				
	g to International Patent Classification (IPC) or to both f			
	: CO7K 7/40; COIN 33/532, 33 / 533			
	L: 435/964,968; 436/543,544,546,547,800	,819,822,823; 530/303		
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	"PHOSPHONAT?"			
III. DOCI	UMENTS CONSIDERED TO BE RELEVANT			
Caledota ,	Citation of Document, 11 with indication, where a	ppropriate, of the relevant passages 17	i Relevant to Claim No. 13	
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	US,3, 4,014,861 (Geiger e	t 51 \ 20 March		
$\frac{X}{Y}$	1977, see column 2, formu		$\frac{12-22}{1-11,23-46}$	
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	11708 7-11 and 56.		!	
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	column 3, line 16; claim	i •	:	
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Y	1'S, 1, 4.610, 868 (Fountain)		1-46	
	September 1986, see colum	n 4, lines 47-50.		
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CO	naidered to be of particular relevance	invention	or theory underlying the	
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"L" document which may throw doubts on priority claim(s) or involve an inventive step				
which is cited to establish the publication date of another citation or other special reason (as specified) "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the				
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. "P" document published prior to the international filing date but in the art.				
later than the priority date claimed "&" document member of the same patent family				
IV. CERTIFICATION				
Date of the Actual Completion of the International Search Date of Malling of this International Search Report				
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International Searching Authority Signature of Authorized Officer Mary E. CEPERLEY MARY E. CEPERLEY				
TSA/IE		MARY E. CEPERLEY	- 1	